This Page Is Inserted by IFW Operations and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

As rescanning documents will not correct images, please do not report the images to the Image Problem Mailbox.



X06936	ubulln/Myosin			
700330	alpha-tubulin ·	24h, 6X	U	3
R59199	est = X79535 - beta tubulin	24h, 4X	Ū	3
224727	tropomyosin isoform	24h, 4X	D	3
T92451	est = M12125 - tropomyosia	24h, 5X	D	3
X05276	fibroblast tropomyosin TM30 (pl)	24h, 4X	D	3
T60155	est = JO5192 - alpha-actin (ACTA)	24h, IIX	D	3
M19283	Kamma-actin	24h, 4X	D	3
X54163	Cardiac troponin I	24h, 5X	D	3
H44011	myosin heavy chain	8h, 5X	D	3
T55741	est = X85337 - myosin light chain kinase (MLCK)	24h, 6X	D	3
X53416	actin-binding protein (filamin) (ABP-280)	24h, 5X	D	
T97948	neutral calponin	24h, 15X	D	3
Cell Cy	cle	· · · · · · · · · · · · · · · · · · ·		
L49231	retinoblastoma susceptibility protein (RBI)	24h, 12X	Ū	3
X39798	PRAD1 mRNA for cyclin	24h, 9X	D	3
D13891	M-2H	8h.24h, 6-7X	ō	3
L13698	gas-1 (growth-arrest-specific gene)	8h,24, 6-15X	Ď	i
X62048	Weel hu	8h.24h, 4-5X	Ď	3
H50438	est = M81934 - ede25B	8h, 4X	6	3
U09477	53BPI p53-binding protein	Bh, SX	ŏ	3
Cosgula		1 84 37	10	
R16659	est = L27624 and AC002076 - TFPI-2	24h, 8X	U	1
M59499	lipoprotein-associated coagulation inhibitor	8h, 6X	Б	3
M14083	beta-migrating plasminogen activator inhibitor i	24h. 4X	Б	3
102933	blood coagulation factor VII gene	24h. 5X		3
Comple		14714 7/	D	رون
M31516		8,24h, 4-5X		-
T54547	est = M84526 - adipsin/complement factor D		ν _σ	Ļ
T69603	-cst-=-M14058 complement-cir-component	24h, 7X 24h, 4X	D	3
R12676	est = M65292 - 1.4-kb mRNA of complement factor H		D	
	es/Receptors	40', 4X	D	3
MZTTZT	nnles	a. 79		
M29150		8h, 6X	Ų.	2
X58377	Imericulin-11 (IL-11)	8h, 10X	U	3
M29696		8,24h, 4-11X	U	1
U02020	Interleulin-7 (IL-7) receptor, alpha chain	24h, 4X	U	1
T61446	pre-B. cell enhancing factor (PBEF) A20	8h,24h, 4X	U	3
		8h, 4X	U	.3
L35263	CSaids binding protein (CSBP1)	8h, 7X	D.	3
M58286	tumor necrosis factor receptor	8h,24h, 4-5X	D	1
H14506	est = L36034 - pre-B cell mimulating factor	24h, 4X	D	1
X72012	endoglin	8h, 23X	D	3
	ular Matrix and Cell Adhesion		·	
L36531	Integrin alpha 8 subunit	40', 6X	U	3
M80744	E16, an integral membrane protein	40', 4X	U	3
X14787	thrombospondin 1 (TSP1)	24h, 21X	D	1
112260		24ኪ 13X	_	
L12350	thrombospondin 2 (TSP2)		D	3
X05231	collagenase	40',8h, 4-7X	D	3
X05231 L25285	collagenase collagen alpha-I type XV (COLISAI)	40',8h, 4-7X 24h, 13X	D D	3
X05231 L25285 J03464	collagenase collagen alpha-I type XV (COLISAI) collagen alpha-2 type I	40',8h, 4-7X 24h, 13X 24h, 14X		3
X05231 L25285 J03464 M11718	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X		3
X05231 L25285 J03464 M11718 M34570	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type V collagen alpha-2 type VI	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X	وامامام	3 3 3
X05231 L25285 J03464 M11718 M34570 T51558	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI ext = K01228 - procollagen alpha I(I) chain	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 14X 24h, 8X 24h, 6X 24h, 6X		3 3 3 3
X05231 L25785 J03464 M11718 M34570 T51558 X06700	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI ext = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen	40,8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 6X	وامامام	3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI cotlagen alpha-2 type VI ext = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen bys/1 oxidase, an extracellular copper enzyme	40,8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X	ووووووو	3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type Vi est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyxyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprotein	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 4X 40',8,24h,5-7X		3 3 3 3 1 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen lyryl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprocein gamma-tarcoglycan, a dystrophia-associated protein	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 4X 40',8,24h,5-7X 24h, 6X	ووووووو	3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gramma-sarcoglycan, a dystrophia-associated protein Tissue labibitor of metalloproteinases-3 (TIMP3)	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 4X 40',8,24h,5-7X		3 3 3 3 1 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI collagen alpha-2 type VI cst = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lytyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gramma-sarcoglycan, a dystrophia-associated protein Tissue labibitor of metalloproteinsses-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 4X 40',8,24h,5-7X 24h, 6X		3 3 3 3 1 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyxyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gamma-aarcoglycan, a dystrophin-associated protein Tissue labibitor of metalloproteinases-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5)	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 4X 40',8,24h,5-7X 24h, 6X 24h, 4X 24h, 42 40',8,24h,5-7X 24h, 6-8X	وووو وووووو	3 3 3 3 1 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gamma-asrooglycan, a dystrophin-associated protein Tissue lahibitor of metalloproteinase-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) casherin-II	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 5-7X 24h, 6X 8h,24h, 5-9X		3 3 3 3 1 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI collagen alpha-2 type VI cst = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen lytyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoproccin gramma-sarcoglycan, a dystrophia-associated protein Tissue labibitor of metalloproteinase-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-II dding Proteins	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 6X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 3-9X		3 3 3 3 1 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-tarcoglycan, a dystrophin-associated protein framma-tarcoglycan, a dystrophin-a	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 6X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 3-9X		3 3 3 3 1 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP B16 H67367 T93295	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyryl oxidase, an extracellular copper enzyme fibolin-I C - a secreted glycoprotein gamma-sarcoglycan, a dystrophin-associated protein Tissue lahibitor of metalloproteinase-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-II dding Proteins est = D38076 - RanBPI (Ran-binding protein I) estihomo of L20294-mouse GTP-binding protein	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 4X 40',8,24h,5-7X 24h, 6S 24h, 6S 24h, 5-9X 24h, 4X 24h, 4X 24h, 4X		3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M1718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 W34976 U14394 U34976 U14394 U34976 U163877 L34056 GTP Bit H67367 T93295 X75593	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI collagen alpha-1 type VI collagen alpha-2 type VI collagen alpha-1 type VI collagen alpha-2 type VI collagen alpha-2 type VI co	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 5-9X 24h, 5X		3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 U03877 U34976 GTP Bit H67367 T93295 X75593 R53966	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI cst = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyryl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gramma-sarcoglycan, a dystrophin-associated protein Tissue labibitor of metalloproteinsses-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cstdherin-II dding Proteins cst = D38076 - RanBP1 (Ran-binding protein I) esti-homo, of L20294-mouse GTP-binding protein ab 13 est = Z22641 - a2-chimacria	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 6X 24h, 5-9X 24h, 5-9X 24h, 3-9X 24h, 5-9X 24h, 5-8X		3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X33743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T93295 X75593 X75593 R53966 H19201	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen lysyl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-tarcoglycan, a dystrophin-associated protein framma-tarcoglycan, a dystrophin-	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 5-9X 24h, 5-9X 24h, 5-9X 24h, 5X 24h, 5X 24h, 5X		3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T93295 X75593 R53966 H19201 Heat Sh	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyryl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-sarcoglycan, a dystrophin-associated protein Tissue lahibitor of metalloproteinases-3 (TIMP3) cat = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-I1 dding Proteins est = D38076 - RanBP1 (Ran-binding protein I) est: borno, of L20294-mouse GTP-binding protein rab 13 est = Z22641 - a2-chimaeria est: borno, to mouse L07974 - gua nucl. disso, stim. ock and Stress-Inducible Proteins	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6-8X 8h, 24h, 5-9X 24h, 4X 24h, 9X 24h, 9X 24h, 9X 24h, 9X		3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T793295 X75593 R53966 H19201 Heat Sh R08183	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen lysyl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-tarcoglycan, a dystrophin-associated protein framma-tarcoglycan, a dystrophin-	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 5-9X 24h, 5-9X 24h, 5-9X 24h, 5X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T793295 X75593 R53966 H19201 Heat Sh R08183 T66307	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyryl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-sarcoglycan, a dystrophin-associated protein Tissue lahibitor of metalloproteinases-3 (TIMP3) cat = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-I1 dding Proteins est = D38076 - RanBP1 (Ran-binding protein I) est: borno, of L20294-mouse GTP-binding protein rab 13 est = Z22641 - a2-chimaeria est: borno, to mouse L07974 - gua nucl. disso, stim. ock and Stress-Inducible Proteins	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 8X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5X 24h, 6-8X 8h,24h, 5-9X 24h, 4X 24h, 5X 24h, 5X 24h, 4X 24h, 4X 24h, 9X 24h, 8X 8h,9X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T793295 X75593 R53966 H19201 Heat Sh R08183	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoprocein gamma-sarcoglycan, a dystrophia-associated protein Tissue labibitor of metalloproteinase-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) esdherin-II dding Proteins est = D38076 - RanBPI (Ran-binding protein I) est homo. of L20294-mouse GTP-binding protein ab 13 est = Z22641 - a2-chlmaeria est: homo. to mouse L07974 - gua. nucl. disso. stim. ock and Stress-Inducible Proteins est = X78821, U07550 - chaperonin 10 heat shock 70xDa protein I	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 4X 24h, 9X 24h, 9X 24h, 9X 24h, 8X 24h, 9X 24h, 8X 24h, 9X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T793295 X75593 R53966 H19201 Heat Sh R08183 T66307	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cst = K01228 - procollagen alpha I(I) chain pro-alpha I(III) collagen lyryl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprocein gamma-sarcoglycan, a dystrophin-associated protein fistrue labibitor of metalloproteinases-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-11 dding Proteins cst = D38076 - RanBPI (Ran-binding protein I) cst:homo. of L20294-mouse GTP-binding protein rab 13 cst = 272641 - a2-chimaeria cst: homo. to mouse L07924 - gua auct. disso, stim. ock and Stress-Inducible Proteins cst = X75821, U07550 - chaperonin 10 heat shock 70xDa protein I transformation-sensitive protein	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 5-9X 24h, 6X 24h, 5-9X 24h, 4X 24h, 9X 24h, 5X 24h, 4X 24h, 9X 24h, 8X 8h, 9X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X33743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T93295 X75593 X75593 R53966 H19201 Hest Sh R08183 T66307 M86752	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lyryl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprotein gamma-tarcoglycan, a dystrophin-associated protein Tissue labibitor of metalloproteinases-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-11 dding Proteins cst = D38076 - RanBPI (Ran-binding protein I) est:homo. of L20294-mouse GTP-binding protein rab 13 est = Z22641 - a2-chimacria est: borno. to mouse L07924 - gua nucl. disso. stim. cek and Stress-Inducible Proteins est = X75821, U07550 - chaperonin 10 heat shock 70xDa protein I transformation-sensitive protein heat shock protein, E coli Dnal homologue	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 16X 24h, 5-7X 24h, 6X 24h, 5-7X 24h, 6X 24h, 5-9X 24h, 5-9X 24h, 4X 24h, 9X 24h, 8X 8h, 9X 24h, 8X 24h, 8X 8h, 4X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 T64394 H67367 T93295 X75593 R53966 R08183 T66307 M86752 L08069	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprocein gamma-arroglycan, a dystrophin-associated protein Tissue labibitor of metalloproteinase-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) catherin-11 eding Proteins est = D38076 - RanBP1 (Ran-binding protein 1) est:homo. of L20294-mouse GTP-binding protein rab 13 est = Z22641 - a2-chimaeria est: homo. to mouse L07924 - gua. nucl. disso. stim. ock and Stress-Inducible Proteins est = X75821, U07550 - chaperonin 10 heat shock 70kDa protein I transformation-sensitive protein best shock protein, E. coli Dnal homologue est = M14328 alpha enolase	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 5X 24h, 5-7X 24h, 5-7X 24h, 5-9X 24h, 4X 24h, 5X 24h, 4X 24h, 5X 24h, 4X 24h, 5X 24h, 5X 24h, 5X 24h, 5X 24h, 8X 8h, 9X 24h, 8X 8h, 9X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X56700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T93295 X75593 R53966 H19201 Heat Sh R08183 T66307 M86752 L08069 H55758 T93272	collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type VI collagen alpha-2 type VI cot = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lytyl oxidase, an extracellular copper enzyme fibulin-I C - a secreted glycoproccin gramma-sarcoglycan, a dystrophia-associated protein fissue labibitor of metalloproteinase-3 (TIMP3) cst = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) cadherin-II dding Proteins est = D38076 - RanBPI (Ran-binding protein I) estihomo. of L20294-mouse GTP-binding protein sb 13 est = Z22641 - a2-chlmacria est: horno, to mouse L07924 - gua nucl. disso, stim. ock and Stress-Inducible Proteins est = X75821, U07550 - chaperonin I0 heat shock 70kDa protein I transformation-sensitive protein heat shock protein, E coli Dnal homologue est = M62829 early growth response protein I	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 5-7X 24h, 6X 24h, 6-8X 8h, 24h, 5-9X 24h, 6-8X 8h, 24h, 5-9X 24h, 4X 24h, 9X 24h, 9X 24h, 8X 8h, 9X 24h, 8X 24h, 11X 24h, 8X 8h, 4X 24h, 15X 824h, 15X 824h, 15X 824h, 15X 824h, 15X 824h, 5-6X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
X05231 L25285 J03464 M11718 M34570 T51558 X06700 L16895 X53743 U34976 U14394 R32771 U03877 L34056 GTP Bit H67367 T93295 X75593 R53966 H19201 Heat Sh R08183 T66307 M86752 L08069 H55758	collagenase collagen alpha-1 type XV (COLISAI) collagen alpha-2 type I collagen alpha-2 type V collagen alpha-2 type V collagen alpha-2 type VI est = K01228 - procollagen alpha I(I) chain pro-alpha I(II) collagen lysyl oxidase, an extracellular copper enzyme fibulin-1 C - a secreted glycoprocein gamma-arroglycan, a dystrophin-associated protein Tissue labibitor of metalloproteinase-3 (TIMP3) est = U37791 - rasi-1 matrix metalloproteinase extracellular protein (S1-5) catherin-11 eding Proteins est = D38076 - RanBP1 (Ran-binding protein 1) est:homo. of L20294-mouse GTP-binding protein rab 13 est = Z22641 - a2-chimaeria est: homo. to mouse L07924 - gua. nucl. disso. stim. ock and Stress-Inducible Proteins est = X75821, U07550 - chaperonin 10 heat shock 70kDa protein I transformation-sensitive protein best shock protein, E. coli Dnal homologue est = M14328 alpha enolase	40',8h, 4-7X 24h, 13X 24h, 14X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 16X 24h, 5X 24h, 5-7X 24h, 5-7X 24h, 5-9X 24h, 4X 24h, 5X 24h, 4X 24h, 5X 24h, 4X 24h, 5X 24h, 5X 24h, 5X 24h, 5X 24h, 8X 8h, 9X 24h, 8X 8h, 9X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3



7			
Interf M2459		T	_
M8743		8h,24h, 5-19X 8h, 6X	10
X0287	(2'-5') oligo A synthetase E (1.8 kb RNA)	8h,24h, 6-15X	۱ü
X0287	(2'-5') oligo A synthetase E (1.6 kb RNA)	8h, 8X	Τŭ
M8728		8h, 8X	Ū
L03072		8h,h, 34X	U
X1594		24h, 7X	ļ۷
X6732	5 Interferon-alpha inducible gene; p27 gene	8h, 5-12X 24h, 7-28X	1 <u>4</u>
H05300	Interferon-Induced guanylate-binding protein I	8h,24h, 9-27X	Ηü
M5554	2 guanylate binding protein isoform II	8h, 9X	۲ŭ
D3188		8h,24h, 3-4X	Ťů
X82200		24h, 14X	Ū
X0249		8h,24h, 4-5X	Ų
M1375		8h,24h, 3-6X	ŲΨ
M2862		8,24h, 55-260X 8h,24h, 10-31X	U
X17668	Indoleamine 2,3-dioxygenase	8h,24h, 9-140X	
M3388		8h,24h, 5-118X	
M3081		8h,24h, 23-53X	Ū
X56841		8h,24h, 5-19X	U
M6061		24h, 32X	Į Ų
M7377		8h,24h, 4-16X 8h,24h, 3-6X	Ų
		1811,2411, 3-07	Ţυ
Kinase R39857	/Phosphatase		
H02889	est = X97630 - serine/threonine protein kinase EMK est = Y11366 IMPA gene	24h, 4X	Ū
U25994		24h. 8X	บั
D21209		Bh, 4X	Ū
X77278	HYL tyrosine kinase	8h,24h, 4-6X 24h, 8X	D
R60908	est = X74764 - receptor protein tyrosine kinase	24h, 21X	<u>D</u>
H65441	est = U78027, L35265 - Bruton's tyrosine kinase	24h, 7X	<u>P</u>
Lamins	proto-oncogene tyrosine-protein kinase (abl)	24h, 4X	<u>D</u>
	laminin B2 chain		<u>~</u>
X79683	bata2 laminin	24h, 4-8X	D
R43734	S78569 - Isminin alpha 4 chain	1 / 4 h 4 v	D
T55218	est = M13452 or M13451 - Jamin A or C	Rh 269	<u>D</u>
Ligands L13740			<u>D</u>
U12767	TR3 orphan receptor mitogen Induced nuclear orphan receptor	8.24h.6-46X	Ū
M77140	pro-galania	Bh. SX	Ū
M63888	heparin-binding growth factor receptor		U
T70920	est = M88279 - Immunophilin (FKBP52)	341 834	<u>v</u>
M20132 M19481	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	PL AV	<u>D</u>
T71662	follistatin est = M14118 - Insulin-like growth factor (IGF-II)	24h. 6-15X	Ď.
M35878	Insulin-like growth factor-binding protein-3		D
M65062	Insulin-like growth factor-binding protein-5		<u>D</u>
M35410	Insulin-like growth factor binding protein 2	400 00 00 00	<u> </u>
X04434	Insulin-like growth factor 1 receptor.	400 00 000	0
L07594 M21574	transforming growth factor-beta type III receptor	8h,24h, 4-6X	5
H88938	platelet-derived growth factor receptor alpha est = M60828 M25295 - keratinocyte growth factor	8h,24h, 12-25X	5
X62381	activin receptor, growth factor-like receptor		<u> </u>
R45296	est = U67784 - orphan G protein-coupled receptor	2/1 /2	2
X02157	fetal erythropoictia		5
M64497	apolipoprotein Al regulatory protein (ARP-1)	1 0) 4 434	ź
L00352	17 beta hydroxysteroid dehydrogenase type 2	24h, 7X	5
M10065	low density lipoprotein (LDL) receptor apolipoprotein E		Σ
Prostagi		40'.8h, 7-26X	2
U04636	cyclooxygenase-2 (cox2)	8h, 7X	,
M72393	cytosotic phospholipase A2 (cPLA2)	8h,24h, 7-12X	
X05908	lipocortin	24h, 9X [_
D38145 Protein	prostacyclio synthase Degradation	24h, 4X	_
T56244	est = D26599 - proteasome subunit HsC7-1	1577 20	-
T54276	proteasome pubunit LMP7 (alkle LMP7C)	24h, 6X L	
T92259	proteasome component IOTA chain	8h, 4X L 24h, 5X L	
D00762	proteasome component HC8	24h, 6X	
D00760	proteasome component C3	24h, 5X U	
H05893	26S protessome subunit p97	24h, 4X	_
M91670	26S protesse (S4) regulatory subunit. ubiquitin carrier protein (E2-EPF)	24h, 16X U	
R67921	est = D55696 - putative cysteine protesse	24h, 4X U 8h, 4X U	
T56256	est = U20657 - ubiquitin protesse proto-oncogene	8h, 4X D	
303589	uoiquitin-like protein (GdX)	24h, 7X D	
T50500	est = 222658 - a placental thrombin inhibitor (PTD)	24h 4X D	



	-oncogenes .			
X8998	5 BCL7B gene	8h,24h, 5-10X	Т	Ū
H4812	The state of the s	8h, 4X		Ŭ
M837	l arginine-rich protein (ARP)	24h, 11X		Ū
M2790	The state of the s	24h, 16X	1	Ū
T5313		8h, 4X	7	D
X1641		24h, 4X	1	D
D4396		8h,24h, 4X	7	D
X8220		8h,24h, 4-7X	7	D
Spilcin				
X13482	U2 small noclear ribonucleoprotein A'	24h, 4X	T	ī
M1584	U2 small suclear RNA-associated B*	24h, 4X	Ti	ī
R41349	splicing factor, arginine/serine-rich 7 (SFRS7)	24h, 10X	Ti	ī
Transc	ription Factors			-
R55041	est = 103161 - serum response factor (SRF)	24h, 5X	π	ī
H46624	est: homology to mouse AB012276 - ATFx	8h, 4X	π	ĵ.
M97676	homeobox protein (HOX7)	8h, 4X	T	Ţ
R26139	est = X59268 - TFIIB	8h, 8X	T	5
M69043	MAD-3, encodes an I kappa B-like protein	8h,24h, 6-10X	T	ī
R26146	NF-kB p105 subunit	24h, 4X	T	;
U09825	acid finger protein (AFP)	8h,24h, 5-8X	Ţ	Ī
H88261		8h, 4X	L	Ī
X03348	beta-glucocorticoid receptor	8h,24h, 4X	Ū	2
M83667	NF-IL6-beta protein	8h,24, 4-6X	I	2
X57435	transcription factor AP-4	8h,24, 4X	Ţ	
M36711	sequence-specific DNA-binding protein (AP-2)	8h, 6X	Ţ	
Y00345	polyA blading protein	24h, 5X	I	
R39209	est = X65644 - MBP-2 for MHC binding protein 2	24h, 5X][_
L19872	aryl hydrocarbon receptor (AhR)	8h,24h, 4-8X	10	?
Transta	tion Factors	7		
R72300		24h, 149X	Ţυ	
H04333	est = U49436 - translation initiation factor 5 (elfs)	74h 4X	Ju	
T69446	est = D13748 - Initiation factor 4A-1 (elF-4A1)	24h, 36X	Ţυ	
L18960	translation initiation factor (cIF-4C)	24h, 39X	Ţ	
H01943	eukaryotic initiation factor 4E (eIF-4E)	74h 4X	Ţυ	
X 62570	IFP53, trypotophanyl-tRNA synthetase	8h,24h, 6-800X	Ţυ	
M81592	gamma-glutamyl carboxylase (hGC)	24h, 5X	13	
212830	SSR (signal-sequence receptor) alpha subunit	24h, 5X	Ţυ	
R60357	alanyi tRNA synthetase	Bh. 5X	LD	
D31762	KJAA0057 gene - homolog of TRAMP	24, 8X	Ιo	?
Miscell	ancous			
M34551		8h.24h, 4-12X		
U33286	chempeone teameries I i our		<u>U</u>	
	Tentomosonic segregation gene homolog CAS		Τū	
T88721	chromosome segregation gene homolog CAS est = U52100, X94770 - XMP mRNA	24h, 4X 24h, 4X	l L	
M26697	est = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23	24h, 4X 24h, 4X	UUU	
	est = U52100, X94770 - XMP mRNA	24h, 4X	l L	
M26697	est = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23	24h, 4X 24h, 4X 24h, 12X	UUU	
M26697 J05682	est = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vas C)	24h, 4X 24h, 4X 24h, 12X	UUU	
M26697 J05682 M22349	cit = U52100, X94770 - XMP mRNA nuckolat phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase	24h, 4X 24h, 4X 24h, 12X	UUU	
M26697 J05682 M22349 M24470	ett – U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X	טט	
M26697 J05682 M22349 M24470 T47964	ett = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine muckoside phosphopolase	24h, 4X 24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X	000	13
M26697 J05682 M22349 M24470 T47964 U07681	ett = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase plocose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = nubunit	24h, 4X 24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X	שטטט	3 1 1 3
M22349 M22349 M24470 T47964 U07681 R42235	cti = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP2C ene for cytochrome PASO	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 8X 24h, 6X	טטטט ט	3 3 1 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120	eti = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lytosomal slaloglycoprotein	24h, 4X 24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X 24h, 9X 24h, 6X 24h, 6X	00000	3 3 1 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123	est = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine suckoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene	24h, 4X 24h, 12X 24h, 12X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X 24h, 6X 24h, 6X 24h, 6X		3 3 1 3 1 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131	cit = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPasc (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckoside phosphorylase NAD(H)-specific isodirate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 6X 24h, 4X 24h, 8X 24h, 8X 24h, 8X 24h, 8X		3 1 3 1 3 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244	cti = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructoklnase est = X57522 - antigen pexide transporter I	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 8X 24h, 6X 24h, 4X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X		3 3 1 3 3 3 3
M22349 M24470 T47964 W07681 R42235 H53120 J00123 H28131 R42244 R26294	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vst C) neuron-specific gamma-2 enolase glucose-6-phosphoste dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X54842 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64184 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids	24h, 4X 24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 9X 24h, 8X 24h, 8X 24h, 8X 24h, 20X 24h, 20X 24h, 4X		3 3 1 3 3 3 3 3
M22349 M24470 T47964 R42235 H53120 J00123 H28131 R42244 R26294 L25270	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine aucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169	24h, 4X 24h, 4Z 24h, 12X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 9X 24h, 6X 24h, 9X 24h, 8X 24h, 8X 24h, 20X 24h, 6X 24h, 4X 24h, 8X 24h, 8X 24h, 4X 24h, 8X		3 3 1 3 3 3 3 3 3
M26697 J05682 M22349 M24490 T44964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833	cti = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPasc (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckoside phosphorylase NAD(H)-specific isodirate dehydrogenase « subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X		3 3 1 3 3 3 3 3 1
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850	cit = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckoside phosphorylase NAD(H)-specific isodirate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofnetoklnase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 s putative nuckotide-binding protein est = X56351 5 - aminofevulinate synthase	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X 24h, 6X 24h, 4X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X		3 3 1 3 3 3 3 3 1
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphoste dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2)	24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X 24h, 9X 24h, 8X 24h, 8X 24h, 20X 24h, 20X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h,24h, 4X		3 1 3 3 3 3 3 3 3 1
M22349 M22349 M2470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 L25270 U01833 H72850 U29195 U19523	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose 6-phosphote dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP2C pene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = APD17732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X563515 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I	24h, 4X 24h, 12X 24h, 12X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 9X 24h, 8X 24h, 8X 24h, 20X 24h, 20X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X		3 3 3 3 3 3 3 3 1
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980	cti = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPasc (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckoside phosphorylase NAD(H)-specific isodirate dehydrogenase « subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M6484 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nuckoside-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentarxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 8X 24h, 6X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 4X 24h, 5X		3 3 1 3 3 3 3 1 3 1 3 3
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleotide phosphorylase NAD(H)-specific Isocitrate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platekt 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2	24h, 4X 24h, 12X 24h, 12X 24h, 12X 24h, 12X 24h, 12X 24h, 12X 24h, 14X 24h, 14X		3 3 1 3 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3 3
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vst C) neuron-specific gamma-2 enolase glucose-6-phosphoste dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X54842 - CYP2C pene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64184 - platelet 6-phosphofructokinase est = X57352 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclobydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 9X 24h, 9X 24h, 8X 24h, 8X 24h, 20X 24h, 4X 24h, 20X 24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 4X 24h, 5X 8h,24h, 4X		3 3 1 3 3 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3
M22349 M22349 M24470 T41964 U07681 R42235 H53120 J00123 H28131 R42244 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X69978	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isodirate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene ett = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nucleoside-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF ERCCS excision repair protein	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 20X 24h, 6X 24h, 4X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 24h, 5X 24h, 6X 24h, 6X 24h, 6X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 24h, 5X 8h, 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X		3 3 1 3 3 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3
M26697 J05682 M22349 M24470 T44964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X69978	cit = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckoside phosphorylase NAD(H)-specific Isodirate dehydrogenase = subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofnuctoklnase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nuckotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF ERCC5 excision repair protein acrodema pigmentosum group C compil factor	24h, 4X 24h, 6X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 8X 24h, 8X 24h, 6X 24h, 4X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 5X 8h,24h, 4X 24h, 4X		3 3 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3 3 3 3
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X65024 R60318	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphosphorylase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56432 - CYP7C gene for cytochrome P450 cst = D12676 - lysosomal slaloglycoprotein enkephalin gene cst = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter I cst = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF ERCCS excision repair protein zeroderma plementosum group C compl. factor est = AF022813 - tetrastran (NAG-2)	24h, 4X 24h, 12X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 8h, 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 6-7X 24h, 4-8X 8h, 24h, 8-7X 24h, 8-7X 8h, 24h, 6-7X 24h, 4-8X 8h, 24h, 8-7X 24h, 4-8X 8h, 24h, 4-8X 8h, 24h, 4-8X 8h, 24h, 4-8X 24h, 4-8X		3 3 1 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3 3 3
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 L25270 U01833 H72850 U29195 U19523 H03980 L19779 L29678 X69978 X69978 X65024 R60318 H87106	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vst C) neuron-specific gamma-2 enolase glucose-6-phosphoste dehydrogenase purine nucleoside phosphorylase NAD(H)-specific leocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - Jysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen pexide transporter I est = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitf ERCC5 excision repair protein zeroderma pigmentosum group C compl. factor est = AF023906 - T245 protein (T245)	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 9X 24h, 8X 24h, 8X 24h, 20X 24h, 4X 24h, 1X 24h, 2X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 8h, 24h, 6-7X 24h, 48X 24h, 48X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
M22349 M24470 M24470 M24470 M24470 M24470 M24470 M247964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U19523 H03980 L19779 Z29678 X69978 X69978 X65024 R60318 H87106 U14650	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPase (vat C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isodirate dehydrogenase « subunit est = X56452 - CYPZC gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructokinase est = X57522 - antigen peptide transporter i est = AF017732 - chromosome 2 cosmids XE169 a putative nucleoside-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF ERCCS excision repair protein zeroderma plementosum group C compl. factor est = AF022813 - tetraspan (NAG-2) est = AF043906 - T245 protein (T245) platelet-endothelial tetraspan nitigen 3	24h, 4X 24h, 6X 24h, 6X 8h,24h, 4-14X 24h, 8X 24h, 8X 24h, 8X 24h, 6X 24h, 6X 24h, 6X 24h, 6X 24h, 4X 24h, 20X 24h, 6X 24h, 4X 24h, 7X 8h,24h, 4X 24h, 7X 8h,24h, 6-7X 24h, 5X 8h,24h, 6-7X 24h, 5X 8h,24h, 6-7X 24h, 8X 24h, 7X 8h,24h, 6-7X 24h, 5X 8h,24h, 6-7X 24h, 8X 24h, 5X 8h,24h, 6-7X 24h, 8X 24h, 5X 8h,24h, 6-7X 24h, 48X 24h, 28X 24h, 28X 24h, 28X 24h, 24X 24h, 24X 24h, 24X 24h, 24X 24h, 24X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X65024 R60318 H87106 U14650 L13385	cit = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B73 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckotide phosphorylase NAD(H)-specific Isocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofnetoklnase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 s putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB00106 - glia maturation factor histone H2A.2 mitf ERCC5 excision repair protein zeroderma pigmentosum group C compl. factor est = AF043906 - T245 protein (T245) platelet-endothelial tetraspan antigea 3 Miller-Dicket Bissencephaly motein (LISI)	24h, 4X 24h, 6X 24h, 6X 8h, 24h, 4-14X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 28X 24h, 4X 24h, 28X 24h, 4X 24h, 4X 24h, 28X 24h, 4X 24h, 28X 24h, 4X		3 3 1 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X65024 R60318 H87106 U14650 L13385 L35263	cit = U52100, X94770 - XMP mRNA nucleolar phosphoprotein B23 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphosphorylase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase purine nucleoside phosphorylase NAD(H)-specific isocitrate dehydrogenase = subunit est = X56452 - CYP7C gene for cytochrome P450 cst = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofructoklnase est = X57522 - antigen peptide transporter I cst = AF017732 - chromosome 2 cosmids XE169 a putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB001106 - glia maturation factor histone H2A.2 mitF ERCCS excision repair protein zeroderma pigmentosum group C compl. factor est = AF022813 - tetraspan (NAG-2) est = AF043906 - T245 protein (T245) platelet-endothelial tetraspan antigen 3 Miller-Dicker Eissencephaly protein (LIS1) CSalds binding protein (CSBP1)	24h, 4X 24h, 4X 24h, 12X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 9X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 8X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 23X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 23X 24h, 4X 24h, 57 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 4X 24h, 57 24h, 4X 24h, 57 24h, 4X 24h, 4X 24h, 57 24h, 4X 8h, 7X		3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
M26697 J05682 M22349 M24470 T47964 U07681 R42235 H53120 J00123 H28131 R42244 R26294 L25270 U01833 H72850 U29195 U19523 H03980 L19779 Z29678 X65024 R60318 H87106 U14650 L13385	cit = U52100, X94770 - XMP mRNA nuckolar phosphoprotein B73 subunit C of V-ATPasc (vst C) neuron-specific gamma-2 enolase glucose-6-phosphate dehydrogenase purine nuckotide phosphorylase NAD(H)-specific Isocitrate dehydrogenase = subunit est = X56452 - CYP2C gene for cytochrome P450 est = D12676 - lysosomal slaloglycoprotein enkephalin gene est = M64784 - platelet 6-phosphofnetoklnase est = X57522 - antigen peptide transporter I est = AF017732 - chromosome 2 cosmids XE169 s putative nucleotide-binding protein est = X56351 5 - aminolevulinate synthase neuronal pentraxin II (NPTX2) GTP cyclohydrolase I est = AB00106 - glia maturation factor histone H2A.2 mitf ERCC5 excision repair protein zeroderma pigmentosum group C compl. factor est = AF043906 - T245 protein (T245) platelet-endothelial tetraspan antigea 3 Miller-Dicket Bissencephaly motein (LISI)	24h, 4X 24h, 6X 24h, 6X 8h, 24h, 4-14X 24h, 8X 24h, 8X 24h, 8X 24h, 6X 24h, 6X 24h, 6X 24h, 4X 24h, 20X 24h, 6X 24h, 20X 24h, 6X 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 7X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 5X 8h, 24h, 4X 24h, 4A 8h, 7X 8h, 24h, 4-8X 8h, 24h, 4-8X		3 3 1 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3



X75090	HLA-DR associated protein I (PHAPI)	8h, 4-8X	D	1 3
U08092	histamine N-methyltransferase (HNMT)	8h. 4X	ᡖ	13
D14874	adrenomedullia	8h,24h, 14-18X	6	H
H80262	est = X15422 - mannose-binding protein C		6	1
R88575	est = U55017 M86521 - transketolase (TKT)	24h, 8X	_	
M13994	cytosolic aldehyde dehydrogenase (ALDHI)	24h, 5X	D	1
M22324	aminopeptidase NACD13 mRNA	24h, 5-7X	D	
M12996		24h, 4X	D	3
	glucose-6-phosphate dehydrogenase (G6PD)	24h, 5X	D	1
T52343	est = Y00433 - glutathione peroxidase	24h, 4X	D	13
T64167	est = dihydrodiol dehydrogenase Isozyme DD2	8h,24h, 4-21X	D	13
X58022	corticotropin-releasing factor binding protein	24h; 4X	D	3
U03688	dioxin-inducible cytochrome P450 (CYP1B1)	8h,24h, 4-11X	D	3
H67764	est = USS764 - estrogen sulfotransferase	24h, 5-12X	D	3
D00137	class I alcohol dehydrogenase beta-1 subunit	8h, 5X	D	3
M77836	pyrroline 5-carboxylate reductase (EC 1.5.1.2)	24, 15X	D	3
T57002	est = M29038 - stem cell protein (SCL)	24h, 4X	D	3
D13665	est = D13665 - osteoblast specific factor 2 (OSF-2p1)	24h, 7X	D	3
M22490	bone morphogenetic protein-2B (BMP-2B)	8h, 4X	D	3
M95787	22kDa smooth muscle protein (SM22)	24h, 5X	D	3
Z11559	iron regulatory factor (IRF)	8h,24h, 5X	D	3
T63612	ferritin heavy chain	8h, 4X	D	3
172863	est = M10119 - ferritin light subunit	24h, 287X	D	3
R56881	est = U61166 - Xenopus laevis intersectin homolog	8h, 4-5X	ъ	3
T63052	est: homolog of M. musculusX52102	24h, 5X	Ū	3
D14663	KIAA0107 gene	24h, 4X	Ū	13
R61352	est = D42085 KIAA0095 gene	24h, 6X	Ŭ	3
U16954	AFiq	24h, 5-7X	Ŭ	3
R98335	est = L13744 AF-9 (unknown function)	8h4-5X	ŭ.	-3.
R01072	est: uaknowa	24h, 5X	Ŭ	3
R10664	est; unknown	24h, 5-7X	Ŭ	1
H24245	est: waknown	24h, 4-5X	Ŭ	3
H11036	est: unknown	24h, 6X	Ŭ	3
R42152	est: unknown	24h, 7X	Ŭ	3
R74454	est: unknown	8h.24h. 4X	Ŭ	3
R54359	est: unknown	24h, 10X	ŭ	H
R01124	est: unknown	24h, 15X	Ü	1
R56443	est; unknown		0	3
T70251	est: unknown	24h, 7X	<u> </u>	3
H92205	est; unknown	40',8h, 4X	_	3
T89666	est; unknown	8h, 4X	D	
H13281	est; unknown	8h, 4X	D	3
R60741	est: unknown	24h, 15X	D	3
H16597		8h, 4X	D	3
R40403	est: uaknowa	24h, 4X	D	3
	est: unknown	24h, 6X	D	3
H78404	est: unknown	8h, 4X	D	3
T48692	est: unknown	8h,24h, 6X	D	1
H40677	est: unknown est: homology to mouse AF003234	8h, 6X ,	۵	3
H86071		8h, 5X	D	3